



# SEQUENCE LISTING

<110> Bettiol, Jean-Luc P.  
Thoen, Christiaan AJK

<120> Detergent Compositions Comprising a Mannanase and a  
Soil Release Polymer

<130> Mannanase and soil release polymer

<140> 09/485,650

<141> 2000-02-14

<150> PCT/US98/12027

<151> 1998-06-10

<160> 6

<170> PatentIn Ver. 2.1

<210> 1

<211> 1482

<212> DNA

<213> Bacillus sp.

<400> 1

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<210> 2

<211> 493

<212> PRT

<213> Bacillus sp.

<400> 2

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Ser Thr Gly Phe Tyr Val Asp Gly Asn Thr Leu Tyr Asp Ala Asn Gly  
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Gln Pro Phe Val Met Arg Gly Ile Asn His Gly His Ala Trp Tyr Lys  
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Asp Thr Ala Ser Thr Ala Ile Pro Ala Ile Ala Glu Gln Gly Ala Asn  
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Thr Ile Arg Ile Val Leu Ser Asp Gly Gly Gln Trp Glu Lys Asp Asp  
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Ile Asp Thr Ile Arg Glu Val Ile Glu Leu Ala Glu Gln Asn Lys Met  
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Val Ala Val Val Glu Val His Asp Ala Thr Gly Arg Asp Ser Arg Ser  
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Ile Gly Lys Glu Asp Thr Val Ile Ile Asn Ile Ala Asn Glu Trp Tyr  
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450

455

460

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&lt;211&gt; 1407

&lt;212&gt; DNA

&lt;213&gt; Bacillus sp.

&lt;400&gt; 3

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&lt;212&gt; PRT

&lt;213&gt; Bacillus sp.

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Ser Thr Gly Phe Tyr Val Asp Gly Asn Thr Leu Tyr Asp Ala Asn Gly	35	40	45
Gln Pro Phe Val Met Arg Gly Ile Asn His Gly His Ala Trp Tyr Lys	50	55	60
Asp Thr Ala Ser Thr Ala Ile Pro Ala Ile Ala Glu Gln Gly Ala Asn	65	70	75
Thr Ile Arg Ile Val Leu Ser Asp Gly Gly Gln Trp Glu Lys Asp Asp	85	90	95
Ile Asp Thr Ile Arg Glu Val Ile Glu Leu Ala Glu Gln Asn Lys Met	100	105	110
Val Ala Val Val Glu Val His Asp Ala Thr Gly Arg Asp Ser Arg Ser	115	120	125
Asp Leu Asn Arg Ala Val Asp Tyr Trp Ile Glu Met Lys Asp Ala Leu	130	135	140
Ile Gly Lys Glu Asp Thr Val Ile Ile Asn Ile Ala Asn Glu Trp Tyr	145	150	155
Gly Ser Trp Asp Gly Ser Ala Trp Ala Asp Gly Tyr Ile Asp Val Ile	165	170	175
Pro Lys Leu Arg Asp Ala Gly Leu Thr His Thr Leu Met Val Asp Ala	180	185	190
Ala Gly Trp Gly Gln Tyr Pro Gln Ser Ile His Asp Tyr Gly Gln Asp	195	200	205
Val Phe Asn Ala Asp Pro Leu Lys Asn Thr Met Phe Ser Ile His Met	210	215	220
Tyr Glu Tyr Ala Gly Gly Asp Ala Asn Thr Val Arg Ser Asn Ile Asp	225	230	235
Arg Val Ile Asp Gln Asp Leu Ala Leu Val Ile Gly Glu Phe Gly His	245	250	255
Arg His Thr Asp Gly Asp Val Asp Glu Asp Thr Ile Leu Ser Tyr Ser			

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Thr Glu Trp Asp Tyr Leu Asp	Leu Ser Glu Asp Trp	Ala Gly Gln His
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Leu Thr Asp Trp Gly Asn Arg	Ile Val His Gly Ala Asp	Gly Leu Gln
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Glu Thr Ser Lys Pro Ser Thr	Val Phe Thr Asp Asp	Asn Gly Gly His
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Pro Glu Pro Pro Thr Ala Thr	Thr Leu Tyr Asp Phe	Glu Gly Ser Thr
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Gln Gly Trp His Gly Ser Asn	Val Thr Gly Gly Pro	Trp Ser Val Thr
355	360	365
Glu Trp Gly Ala Ser Gly Asn	Tyr Ser Leu Lys Ala Asp	Val Asn Leu
370	375	380
Thr Ser Asn Ser Ser His Glu	Leu Tyr Ser Glu Gln	Ser Arg Asn Leu
385	390	395
His Gly Tyr Ser Gln Leu Asn	Ala Thr Val Arg His	Ala Asn Trp Gly
405	410	415
Asn Pro Gly Asn Gly Met Asn	Ala Arg Leu Tyr Val	Lys Thr Gly Ser
420	425	430
Asp Tyr Thr Trp His Ser Gly	Pro Phe Thr Arg Ile	Asn Ser Ser Asn
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<211> 1029

<212> DNA

<213> Bacillus sp.

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<211> 362

<212> PRT

<213> Bacillus sp.

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Asn Pro Asn Ala Gln Gln Thr Thr Lys Thr Val Met Asn Trp Leu Ala  
35 40 45

His Leu Pro Asn Arg Thr Glu Asn Arg Val Leu Ser Gly Ala Phe Gly  
50 55 60

Gly Tyr Ser His Asp Thr Phe Ser Met Ala Glu Ala Asp Arg Ile Arg  
65 70 75 80

Ser Ala Thr Gly Gln Ser Pro Ala Ile Tyr Gly Cys Asp Tyr Ala Arg  
85 90 95

Gly Trp Leu Glu Thr Ala Asn Ile Glu Asp Ser Ile Asp Val Ser Cys  
100 105 110

Asn Gly Asp Leu Met Ser Tyr Trp Lys Asn Gly Gly Ile Pro Gln Ile

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Ser Ala Leu Tyr His Asp Ser Trp Thr Leu Asn Lys Gly Glu Ile Trp 340 345 350		
Asn Gly Asp Ser Leu Thr Pro Ile Val Glu 355 360		